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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2009; month=8; day=7; hr=8; min=1; sec=42; ms=458;]

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Reviewer Comments:

<210> 15
<211> 3933
<212> DNA
<213> Pseudomonas sp. HJ-2 (phb locus)

<400> 15
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60

ttcacatccg caaagcgcca gagacttgcc cgctgttcca aggtcttaat taacgaggaa
120

tggттаатгг gtactgcgag caatgcggca cgtatagctc tggtcaccgg tggtatgggc
180

ggtatcggtа cggcgatcag ccagcgctg catcgggatg gcttcaccgt ggtgggtgggc
240

tgtaatccct actccagccg caaggcttcc tggattgcca cgcaactcga ggcgggcttt
300

cattccact gcatcgactg cgacatcacc gactgggata gcacccgcca ggccttcgac
360

atggtgcacg agactgtcgg cccgatcgat gtattggтca аcaatgccgg catcacccgc
420

gacggcactt tccgcaagat gtccccggaa aactggaagg cggtgatcga taccaatctc
480

accggcctgt tcaacacaac caagcaggtc atcgagggca tgctggccaa gggctgggga
540

cgcgatcatca acatctcctc aatcaatggc cagcgaggcc agttcgggca gaccaactac
600

tccgcgggca aggctggcat tcatggcttc agcatggcct tggcccgcga ggtgagtggc
660

aagggcgtga ccgtaataac ggtttccctt ggctacatca agaccgacat gaccgcggcg
720

attcgcccgg acatcctcga agacatgatt actggcattc ccgtggggcg tctcggccag
780

cccgaggaga tcgcctcgat cgtggcctgg ctggcctccg atcagtctgc ctatgccacc
840

ggcgccgact tctcggtgaa tggcggcgatg aacatgcagt gatgcgccat tcgcgcctc
900

gctcagccat gacatgaggt gttccagatg atcgaagtcg ttatcgtcgc cgccactcgc
960

accgccatcg gcgctttcca ggggagcctg gccggcactc ccgccgttga actggggcgcc
1020

acggatgatcc gccgcctgct cgaacagacc gctctggata gcagtcaggt ggatgaagtg
1080

atactcggcc acgtactcac cgccggtgct ggcagaatac cgctcgccag gcancnggtc
1140

Regarding the above <213> response; per 1.823 of the Sequence Rules, the only valid responses are the Genus species of the organism, "Artificial Sequence", or "Unknown". "Artificial Sequence" and "Unknown" require explanation in the <220>-<223> section; please give the source of the genetic material. Please just list the Genus species as the <213> response; put explanatory matter in the <220>-<223> section; please correct all similar sequences.

The n's at locations 608, 1134, and 1136 are not explained above.

<210> 16
<211> 251
<212> PRT
<213> Pseudomonas sp. HJ-2 (NADPH-dependent acetoacetyl-CoA reductase
(phbB))

<400> 16
Met Gly Thr Ala Ser Asn Ala Ala Arg Ile Ala Leu Val Thr Gly Gly
1 5 10 15

Met Gly Gly Ile Gly Thr Ala Ile Ser Gln Arg Leu His Arg Asp Gly
20 25 30

Phe Thr Val Val Val Gly Cys Asn Pro Tyr Ser Ser Arg Lys Ala Ser
35 40 45

Trp Ile Ala Thr Gln Leu Glu Ala Gly Phe His Phe His Cys Ile Asp
50 55 60

Cys Asp Ile Thr Asp Trp Asp Ser Thr Arg Gln Ala Phe Asp Met Val
65 70 75 80

His Glu Thr Val Gly Pro Ile Asp Val Leu Val Asn Asn Ala Gly Ile
85 90 95

Thr Arg Asp Gly Thr Phe Arg Lys Met Ser Pro Glu Asn Trp Lys Ala
100 105 110

Val Ile Asp Thr Asn Leu Thr Gly Leu Phe Asn Thr Thr Lys Gln Val
115 120 125

Ile Glu Gly Met Leu Ala Lys Gly Trp Gly Arg Val Ile Asn Ile Ser
130 135 140

Ser Ile Asn Gly Gln Arg Gly Gln Phe Gly Gln Thr Asn Tyr Ser Ala
145 150 155 160

Xaa Lys Ala Gly Ile His Gly Phe Ser Met Ala Leu Ala Arg Glu Val
165 170 175

Please correct the above <213> response to just indicate the Genus species of the organism; place explanatory matter in the <220>-<223> section. Also, the above <213> response exceeds the Sequence Rules' required 72-character line limit. The "Xaa" at location 161 is not explained above.

<210> 17
<211> 392
<212> PRT
<213> Pseudomonas sp. HJ-2 (beta-ketothiolase (phbA))

<400> 17
Met Ile Glu Val Val Ile Val Ala Ala Thr Arg Thr Ala Ile Gly Ala
1 5 10 15

Phe Gln Gly Ser Leu Ala Gly Thr Pro Ala Val Glu Leu Gly Ala Thr
20 25 30

Val Ile Arg Arg Leu Leu Glu Gln Thr Ala Leu Asp Ser Ser Gln Val
35 40 45

Asp Glu Val Ile Leu Gly His Val Leu Thr Ala Gly Ala Gly Arg Ile
50 55 60

Pro Leu Ala Arg Xaa Xaa Val Ile Ala Gly Leu Pro His Ala Val Pro
65 70 75 80

Please correct the above <213> response. Also, the "Xaa's" at locations 69-70 are not explained above.

Application No: 10583840 Version No: 2.0

Input Set:

Output Set:

Started: 2009-07-22 14:17:12.979
Finished: 2009-07-22 14:17:15.807
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 828 ms
Total Warnings: 18
Total Errors: 6
No. of SeqIDs Defined: 18
Actual SeqID Count: 18

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 402	Undefined organism found in <213> in SEQ ID (12)
W 402	Undefined organism found in <213> in SEQ ID (13)
W 402	Undefined organism found in <213> in SEQ ID (14)
W 402	Undefined organism found in <213> in SEQ ID (15)
E 342	'n' position not defined found at POS: 608 SEQID(15)
E 342	'n' position not defined found at POS: 1134 SEQID(15)
E 342	'n' position not defined found at POS: 1136 SEQID(15)
W 402	Undefined organism found in <213> in SEQ ID (16)
E 341	'Xaa' position not defined SEQID (16) POS (161)

Input Set:

Output Set:

Started: 2009-07-22 14:17:12.979
Finished: 2009-07-22 14:17:15.807
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 828 ms
Total Warnings: 18
Total Errors: 6
No. of SeqIDs Defined: 18
Actual SeqID Count: 18

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (17)
E 341	'Xaa' position not defined SEQID (17) POS (69)
E 341	'Xaa' position not defined SEQID (17) POS (70)
W 402	Undefined organism found in <213> in SEQ ID (18)

<110>	LG CHEM, LTD.	
<120>	Poly(3-hydroxyalkanoate) Block Copolymer Having Shape Memory Effect	
<130>	LC05PCT042	
<140>	10583840	
<141>	2009-07-22	
<150>	KR 10-2005-0059907	
<151>	2005-07-04	
<160>	18	
<170>	KopatentIn 1.71	
<210>	1	
<211>	18	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Choi3 (PCR Primer)	
<400>	1	
	ccgccstgsa tcaagtac	18
<210>	2	
<211>	20	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Choi4 (PCR Primer)	
<400>	2	
	gytsgtgsgygc tcyycgttcc	20
<210>	3	
<211>	24	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	HJ-PHB-N (PCR Primer)	
<400>	3	
	caccatgctg agttgcgctc tagc	24
<210>	4	

<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> HJ-PHB-C (PCR Primer)

<400> 4
tcadmsytty acrtarcgkc ctggygc

27

<210> 5
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> SCL-1 (PCR Primer)

<400> 5
gatcgatacc aatctcaccg

20

<210> 6
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> SCL-2 (PCR Primer)

<400> 6
caaagccagt ggttcgacgt a

21

<210> 7
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> SCL-3 (PCR Primer)

<400> 7
ctgctgaaac tggttgagc

19

<210> 8
<211> 47
<212> DNA
<213> Artificial Sequence

<220>

<223> SD-BA-N (PCR Primer)

<400> 8
gggggtacca ataaggagat atacatatgg gtactgcgag caatgcg 47

<210> 9
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> BA-C (PCR Primer)

<400> 9
cccactagtt cagcgctcga tggccagc 28

<210> 10
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> SD-phbC-N (PCR Primer)

<400> 10
gggcatatga ccagaagaa caacagcg 28

<210> 11
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> phbC-C (PCR Primer)

<400> 11
cccactagtt cadmscttya crtaacgtcc tggcgcygc 39

<210> 12
<211> 756
<212> DNA
<213> Pseudomonas sp. HJ-2

<220>
<221> variation
<222> (482)
<223> n=A, C, G or T

<400> 12

atgggtactg cgagcaatgc ggcacgtata gctctggtca ccggtggtat gggcggtatc	60
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ccctactcca gccgcaaggc ttcttggtatt gccacgcaac tcgaggcggg ctttcacttc	180
cactgcatcg actgcgacat caccgactgg gatagcacc gccaggcctt cgacatggtg	240
cacgagactg tcggcccgat cgatgtattg gtcaacaatg ccggcatcac ccgcgacggc	300
actttccgca agatgtcccc ggaaaactgg aaggcgggtga tcgataccaa tctcaccggc	360
ctgttcaaca caaccaagca ggtcatcgag ggcatgctgg ccaagggtg gggacgcgtc	420
atcaacatct cctcaatcaa tgccagcga ggccagttcg ggcagaccaa ctactccgcg	480
gncaaggctg gcattcatgg cttcagcatg gccttggccc gcgaggtgag tggcaagggc	540
gtgaccgtca atacggtttc ccctggctac atcaagaccg acatgaccgc ggcgattcgc	600
ccggacatcc tcgaagacat gattactggc attcccgtgg gccgtctcgg ccagcccag	660
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<210> 13
 <211> 1179
 <212> DNA
 <213> Pseudomonas sp. HJ-2

<220>
 <221> variation
 <222> (207)
 <223> n=A, C, G or T

<220>
 <221> variation
 <222> (209)
 <223> n=A, C, G or T

<400> 13	
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accgctctgg atagcagtca ggtggatgaa gtgatactcg gccacgtact caccgccggt	180
gctggcagaa taccgctcgc caggcaneng gtcatcgccg gcctgccaca cgccgtaccg	240
gcgatgacc tgaacaaggc ctgtggctcc ggctgaaag ccctgcacct gggcgcccag	300
gccatccgct gtggcgatgc cgagggtggtg attgccggtg gcatggagaa catgagcctg	360
tcgtcctatg tcttgcccaa ggcccgcacc ggctgcgca tgggccacgc gcagctggtc	420

gacagcatga tcgtcgacgg cctgtgggac gccttcaacg actaccacat ggggatcact	480
gccgagaacc tggtagacaa gtacggcatc agccgcgaag cccaggacga attcgccgcc	540
gcctcgcagc agaaagccgt ggccgccatc gagaccggtc gcttccgcga cgagatcgtc	600
cgggtgagca ttccgcagcg caagggcgag gcgtgagct tcgacaccga cgaacagcca	660
cgcgcgggca ccaccgccga gtcgctgggc aagctgaaac cggccttcaa gaacgacggc	720
agcgttactg ccggcaacgc ttccagtctc aacgacggcg ccgccgcggt actgctgatg	780
agtgcggcaa aggccgcagc gcttgggtctg ccagtgtctg cgaagatcgc cgcctacgcc	840
aatgccggcg tcgaccgggc gatcatgggt atcggaccgg tgtcggccac ccgcagttgc	900
ctggagaagg cgggctggag tctggcagag ctggatctga tcgaggccaa tgaagccttc	960
gcggcccagg ccctggccgt gggtcaggag ctgggctggg atgctggcag ggttaacgtc	1020
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agcctgctgc atgaaatgct caggcgcgac gcgaaaaaag gcctcgctac cctgtgtatc	1140
ggtggcgggc agggcggtggc gctggccatc gagcgctga	1179

<210> 14
 <211> 1701
 <212> DNA
 <213> Pseudomonas sp. HJ-2 (SCL-PHA synthase (phaC))

<400> 14	
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ttcgtcctgc agcaactgcg cttatacgtg gcgcaaaata cttggttcag cgggcacgac	120
caaagccagt ggttcgacgt acctgtcgag gcgttggagc aactgcaggc ggactaccaa	180
caacagtggg ccgaacttgg ccagcaattg ctgagctgcc agccgttcgc attcagcgat	240
cgtcgcttcg ccagtggcaa ctggagcgaa ccgctgttcg gttccctggc tgccctctac	300
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ccccgccagc gcttgcggtta cttgatcgag caagcgattg ccgcaagcgc cccaagtaac	420
tttctgctga gcaaccctga tgccctgcaa cgcctagtgg aaaccaggg cgcagccta	480
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ttgggcgatt tcgaagtcgg cgtgaatctg gccaccacce ctggtgccgt ggtactggaa	600
accctctgtt tccagctgat ccagtattcg ccgctcagcg aaacgcaata ccagcggccg	660
atattcatgg tcccgcctg gatcaacaag tactacatcc ttgacctcgg gcccgaaaac	720

tctctaatacc gtcacttact ggagcgaggc catcaagttt ttctgatgtc ctggcgcaac	780
ttcactcagg aacaggccga catcacctgg gagcagatca tccaggacgg agtgatcagc	840
gccctgcgca ctaccggggc catcagtggg gagcgccacc tgaactgttt gggtttctgc	900
atcggcggca ccatgctgag ttgcgtctta gcggtgctgg cagcgcgtgg cgaccaggac	960
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gtcttcgtcg atgagcaact ggtggcctac cgtgagcgca ccatcggtgg ccatggtggc	1080
aaatgtggcc tgttcgcggg tgaggacatg ggcaatacct tctccctgct gcggcccaac	1140
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ggtctactgt tctggaacaa cgacagcacc aatctgccgg ggccctgta ttgctggtat	1260
ctgcgccaca cctacctgca gaacgacctc aaatcggggg agttggatct gtgcggcgtc	1320
aagttggatc tgcggggccat agacgcacca gcctacatct tgggaaccca tgacgaccac	1380
atcgtgccct gggaagcgc ctatgccagc acggaattgc tgggaggtcc aaagcgcttt	1440
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cattactggg tcaatgaaca catagcgccg gtagctgacg actggtgca gggagctcag	1560
cagcattccg gcagttggtg gggtgactgg ttgcctggg tgaccggcta tgccggccca	1620
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ggacgttatg tgaagctatg a	1701

<210> 15
 <211> 3933
 <212> DNA
 <213> Pseudomonas sp. HJ-2 (phb locus)

<400> 15	
gagctcaatg cgcgccagga ctggtgtgcg aggacaaccc ggcgtcaccg ggggacattg	60
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tggttaatgg gtactgcgag caatgcggca cgtatagctc tggtcaccgg tggatgggc	180
ggtatcggta cggcgatcag ccagegcctg catcgggatg gcttcaccgt ggtggtgggc	240
tgtaatccct actccagccg caaggcttcc tggattgcca cgcaactcga ggcgggcttt	300
cacttccact gcacgactg cgacatcacc gactgggata gcacccgcca ggccttcgac	360
atggtgcacg agactgtcgg cccgatcgat gtattgggtca acaatgccgg catcacccgc	420
gacggcactt tccgcaagat gtccccggaa aactggaagg cggtgatcga taccaatctc	480

accggcctgt tcaacacaac caagcaggtc atcgagggca tgctggccaa gggctgggga	540
cgcgatcatca acatctcttc aatcaatggc cagcgaggcc agttcgggca gaccaactac	600
tccgcggnca aggctggcat tcatggcttc agcatggcct tggcccgcga ggtgagtggc	660
aaggggcgtga ccgtcaatac ggtttccctt ggctacatca agaccgacat gaccgcggcg	720
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cccgaggaga tcgcctcgat cgtggcctgg ctggcctccg atcagtctgc ctatgccacc	840
ggcgccgact tctcggtgaa tggcggcatg aacatgcagt gatgcgccat tcgcgccctc	900
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ttcaacgact accacatggg gatcactgcc gagaacctgg tagacaagta cggcatcagc	1440
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accggtcgtc tccgcgacga gatcgtcccg gtgagcattc cgcagcgcaa gggcgaggcg	1560
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ctgaaaccgg ccttcaagaa cgacggcagc gttactgccg gcaacgcttc cagtctcaac	1680
gacggcgccg ccgcggtact gctgatgagt gcggcaaagg ccgcagcgtc tggctctgcca	1740
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ggaccggtgt cggccacccg cagttgcctg gagaaggcgg gctggagtct ggcagagctg	1860
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ggcgctccg gctgccgct actggtcagc ctgctgcatg aaatgctcag gcgcgacgcg	2040
aaaaaaggcc tcgtaccct gtgtatcggg gccggccagg gcgtggcgct ggccatcgag	2100
cgctgagtga cgctttcgcg actctgccgg acgtgcccc ctgcaccgc accgccaggc	2160

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gctgccttct acctgctgaa ttccggtttc ctgctgaaac tgttgagct tctccccatc	2580
gatgagcaga agccccgcca gcgcttgctg tacttgatcg agcaagcgat tgccgcaagc	2640
gccccaaagta actttctgct gagcaaccct gatgccctgc aacgcctagt ggaaaccag	2700
ggcgccagcc tactaagtgg cctgttgcat cttgccagtg acctgcaggc aggcaagttg	2760
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ggagtgatca gcgcctgcg cactaccgg gccatcagtg gtgagcgcca cctgaactgt	3120
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ggcgaccagg acattgccag cctgagtcta ttcgccactt ttcttgacta ccttgatacc	3240
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ggccatggtg gcaaagtgg cctgttcgc ggtgaggaca tgggcaatac cttctccctg	3360
ctgcgcccca acgagctgtg gtggaactac aacgtagaca aatatctcaa ggggcagaag	3420
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tattgctggt atctgcgcca cacctacctg cagaacgacc tcaaatacggg ggagttggat	3540
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catgacgacc acatcgtgcc ctggcgagc gcctatgcca gcacggaatt gctgggaggt	3660
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aggaacaagc gccattactg ggtcaatgaa cacatagcgc cggtagctga cgactggctg	3780
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<210> 16
<211> 251
<212> PRT
<213> Pseudomonas sp. HJ-2 (NADPH-dependent acetoacetyl-CoA reductase (phbB))

<400> 16
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1 5 10 15

Met Gly Gly Ile Gly Thr Ala Ile Ser Gln Arg Leu His Arg Asp Gly
20 25 30

Phe Thr Val Val Val Gly Cys Asn Pro Tyr Ser Ser Arg Lys Ala Ser
35 40 45

Trp Ile Ala Thr Gln Leu Glu Ala Gly Phe His Phe His Cys Ile Asp
50 55 60

Cys Asp Ile Thr Asp Trp Asp Ser Thr Arg Gln Ala Phe Asp Met Val
65 70 75 80

His Glu Thr Val Gly Pro Ile Asp Val Leu Val Asn Asn Ala Gly Ile
85 90 95

Thr Arg Asp Gly Thr Phe Arg Lys Met Ser Pro Glu Asn Trp Lys Ala
100 105 110

Val Ile Asp Thr Asn Leu Thr Gly Leu Phe Asn Thr Thr Lys Gln Val
115 120 125

Ile Glu Gly Met Leu Ala Lys Gly Trp Gly Arg Val Ile Asn Ile Ser
130 135 140

Ser Ile Asn Gly Gln Arg Gly Gln Phe Gly Gln Thr Asn Tyr Ser Ala
145 150 155 160

Xaa Lys Ala Gly Ile His Gly Phe Ser Met Ala Leu Ala Arg Glu Val
165 170 175

Ser Gly Lys Gly Val Thr Val Asn Thr Val Ser Pro Gly Tyr Ile Lys
180 185 190

Thr Asp Met Thr Ala Ala Ile Arg Pro Asp Ile Leu Glu Asp Met Ile
195 200 205

Thr Gly Ile Pro Val Gly Arg Leu Gly Gln Pro Glu Glu Ile Ala Ser
210 215 220

Ile Val Ala Trp Leu Ala Ser Asp Gln Ser Ala Tyr Ala Thr Gly Ala
225 230 235 240

Asp Phe Ser Val Asn Gly Gly Met Asn Met Gln
245 250

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<210>      17
<211>      392
<212>      PRT
<213>      Pseudomonas sp. HJ-2 (beta-ketothiolase (phbA))

<400>      17
Met Ile Glu Val Val Ile Val Ala Ala Thr Arg Thr Ala Ile Gly Ala
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Phe Gln Gly Ser Leu Ala Gly Thr Pro Ala Val Glu Leu Gly Ala Thr
      20              25              30

Val Ile Arg Arg Leu Leu Glu Gln Thr Ala Leu Asp Ser Ser Gln Val
      35              40              45

Asp Glu Val Ile Leu Gly His Val Leu Thr Ala Gly Ala Gly Arg Ile
      50              55              60

Pro Leu Ala Arg Xaa Xaa Val Ile Ala Gly Leu Pro His Ala Val Pro
      65              70              75              80

Ala Met Thr Leu Asn Lys Val Cys Gly Ser Gly Leu Lys Ala Leu His
      85              90              95

Leu Gly Ala Gln Ala Ile Arg Cys Gly Asp Ala Glu Val Val Ile Ala
      100             105             110

Gly Gly Met Glu Asn Met Ser Leu Ser Ser Tyr Val Leu Pro Lys Ala
      115             120             125

Arg Thr Gly Leu Arg Met Gly His Ala Gln Leu Val Asp Ser Met Ile
      130             135             140

Val Asp Gly Leu Trp Asp Ala Phe Asn Asp Tyr His Met Gly Ile Thr
      145             150             155             160

Ala Glu Asn Leu Val Asp Lys Tyr Gly Ile Ser Arg Glu Ala Gln Asp
      165             170             175

Glu Phe Ala Ala Ala Ser Gln Gln Lys Ala Val Ala Ala Ile Glu Thr
      180             185             190

Gly Arg Phe Arg Asp Glu Ile Val Pro Val Ser Ile Pro Gln Arg Lys
      195             200             205

Gly Glu Ala Leu Ser Phe Asp Thr Asp Glu Gln Pro Arg Ala Gly Thr
      210             215             220

Thr Ala Glu Ser Leu Gly Lys Leu Lys Pro Ala Phe Lys Asn Asp Gly
      225             230             235             240

Ser Val Thr Ala Gly Asn Ala Ser Ser Leu Asn Asp Gly Ala Ala Ala
      245             250             255

Val Leu Leu Met Ser Ala Ala Lys Ala Ala Ala Leu Gly Leu Pro Val

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